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GRANT NUMBER: DAMD17-94-J-4411

TITLE: Development of Novel Peptide Inhibitors of the Estrogen Receptor

PRINCIPAL INVESTIGATOR: Richard J. Miksicek, Ph.D.

CONTRACTING ORGANIZATION: State Uni

State University of New York Stony Brook, New York 11794-3366

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13. ABSTRACT (Maximum 200 words)			
This project sime to test th	a hypothesis that nolyn	entides designed to 1	mimic the dimerization domain of
the estrogen recentor are	e hypomesis that poryponable of serving as it	chibitors of receptor	function by virtue of their ability
to interfere with the hindi	ng of receptor protein t	o DNA. During the	e first year of this award, peptides
have been acquired and as	ssavs have been develor	ped to initiate this pi	roject. Several candidate peptides
have been identified that a	ppear to specifically blo	ock the DNA-bindin	g activity of the estrogen receptor
as assessed using an in t	vitro gel mobility shift	t assay. These obs	ervations will be confirmed and
extended during the final	vear of this project by a	issessing whether or	not this inhibitory activity can be
reproduced in a cell cul	ture model system of	estrogen receptor	action. Due to a change in the
institutional affiliation of	the PI (Dr. R.J. Miksice	ek), requests have be	een filed to permit a change in the

14. SUBJECT TERMS

Estrogen Receptor, Breast Cancer, Peptides, Dimerization, Antagonists

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performance site for this project as part of a subcontracting arrangement. This request is currently

pending with the US Army MRDC.

TATELOTICATION 1

#### FOREWORD

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- () In conducting research using animals, the investigator(s) adhered to the "Guide for the Care and Use of Laboratory Animals," prepared by the Committee on Care and Use of Laboratory Animals of the Institute of Laboratory Animal Research Council (NIH Publication No. 86-23, Revised 1985).
- ( ) For the protection of human subjects, the investigator(s) have adhered to policies of applicable Federal Law 32 CFR 219 and 45 CFR 46.
- (X) In conducting research utilizing recombinant DNA technology, the investigator(s) adhered to current guidelines promulgated by the National Institutes of Health.

Principal Trestigator's Signature

Date

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#### DAMD 1794J4411

### INTRODUCTION

Title:

Development of Novel Peptide Inhibitors of the Estrogen Receptor

ID No.: PI: DAMD1794J4411 Richard J. Miksicek

Period:

09/30/94 through 09/29/96

This USAMRDC Breast Cancer Research IDEA Award was made to Richard J. Miksicek, who currently holds appointments as Adjunct Assistant Professor of Pharmacology at The State University of New York at Stony Brook (SUNY @ Stony Brook) and Assistant Professor of Physiology at Michigan State University (MSU). This annual report covers the first year of this award, through 09/29/95.

The aim of this project is to exploit what we have learned through the techniques of modern molecular biology about the functional organization of the estrogen receptor (ER) protein to develop a new class of ER antagonists with a novel mechanism of action. The general aim is to investigate the potential of disrupting subunit dimerization to block the transcriptional stimulatory activity of ER. Essentially all steroid receptor inhibitors developed to date represent steroid analogues that act as hormone antagonists by binding to the ligand-binding site of these proteins. However, instead of stimulating receptor activity as do the physiologically important steroids, these "antihormones" block receptor function through a mechanism that is not entirely understood. As an alternative to hormone antagonists, we are testing the possibility of using peptides to inhibit ER function. These peptides are designed to mimic the dimerization interface of ER and to interfere with subunit association, thereby preventing efficient DNA-binding by receptor. If this approach proves to be feasible, it would have several important advantages over classical hormone antagonists, since they should still be able to inhibit ER function in the face of stimulatory concentrations of estradiol. They should also have a different pattern of resistance than currently available antagonists and they are likely to display very different pharmacokinetic properties. Progress during year one of this award is described in greater detail, below.

During the first year of this award, the P.I. (Richard Miksicek) accepted a new position (effective 07/01/95) in the Department of Physiology at MSU, retaining an appointment as Adjunct Assistant Professor in the Department of Pharmacological Sciences of SUNY @ Stony Brook (letters of appointment included in appendix A). In conjunction with this change, permission was requested from the US Army MRDC to change the site of performance of this project from SUNY @ Stony Brook to MSU by establishing a new subgrant between these institutions (letter of request and a copy of the revised "Description of Facilities and Equipment" are included as appendix B and appendix C). This letter also requested that Dr. Sandra Haslam (Professor of Physiology and Director of the MSU Breast Cancer Program) be named as Principal Investigator for the MSU subgrant. A revised budget outlining an arrangement to subgrant unobligated funds from SUNY @ Stony Brook to MSU has been reviewed and approved by both institutions and has been submitted to the US Army MRDC for approval.

The original statement of work for this project included the following specific tasks:

Task 1, to identify the minimal region of the ER polypeptide able to inhibit the DNA-binding and transcriptional activity of intact ER when co-expressed in cultured cells (months 1-18)

a. to develop a series of plasmids for expressing progressively smaller portions of the ER open reading frame (months 1-6)

b. to perform gel mobility shift assays to ascertain which fragments of ER block DNA-binding when expressed together with intact ER (months 6-12)

c. to confirm that fragments of the ER open reading frame that block DNA-binding activity also function as dominant inhibitors of ER function in vivo (months 12-24)

Task 2, to identify short peptides modeled after the ER dimerization motif that act as inhibitors of DNA-binding and subunit association (months 1-24)

a. to obtain synthetic ER peptides (months 1-3)

b. to test the effect of synthetic peptides on the DNA-binding activity of ER using an *in vitro* gel mobility shift assay (months 3-12)

c. to characterize the inhibitory potency of active peptides and attempt to optimize the effect by

testing additional overlapping peptides (months 6-24)

d. to perform feasibility experiments based on liposome-mediated peptide transfer to assess the ability of selected peptides to inhibit ER function in cultured cells (months 12-24)

## **EXPERIMENTAL METHODS**

Construction of plasmids for efficient expression of ER in cultured mammalian cells.

We have developed a number of cDNA expression plasmids that contain various portions of the protein coding region of the human ER (hER) gene. Several of these plasmids were designed to express a family of variant hER cDNAs have been identified in breast tumors and tumor-derived cell lines (1). Derivatives expressing ERΔE2, ERΔE3, ERΔE5, and ERΔE7 (where the number refers to a deleted exon) are currently available and those expressing ERΔE4 and ERΔE6 will be completed by the end of 1995. These plasmids represent derivatives of pCMV4 (2) and pcDNA3 (InVitrogen Corp., San Diego, CA), which we have shown can support high levels of receptor expression in a variety of mammalian cell lines, including Cos7 (3). In addition to these hER splicing variants, we have a large number of engineered mutants of hER, including a series of carboxy-terminally truncated clones that remove progressively larger portions of hER beginning with domains reported to be important for activation of transcription (4) and subunit dimerization (5). Additional polypeptide expression plasmids will be constructed, as needed, by inserting short (60-180 bp) fragments of the hER cDNA into a vector designed in this laboratory that contains a consensus translation initiation motif, as described in the original application.

Transfection conditions for assessing inhibitory activity of expressed fragments of ER.

To measure the ability of hER polypeptides to interfere with activation of gene expression by intact receptor, transient co-transfection assays are being performed as previously described (6). Briefly, this assay utilizes a calcium phosphate-mediated transfection procedure to introduce plasmids harboring various ER cDNA expression clones into HeLa cells along with an estrogen-responsive reporter plasmid (pERE-TK-CAT). Each transfection experiment includes a plasmid expressing a portion of the hER open reading frame that is transcriptionally inactive on its own, but has the potential of inhibiting co-transfected wild-type receptor. The inhibitory potential of receptor fragments is assessed by measuring the activity of chloramphenicol acetyltransferase (CAT) enzyme produced in cultures of cells transfected with intact receptor alone (pCMV-ER), compared with transfections containing both intact hER and the various fragmentary hER expression clones. Levels of ER expression are being assessed where possible by western blot analysis using an ER-specific monoclonal antibody (Mab-17) that was previously developed in this laboratory (7). For ER polypeptides that retain the Mab-17 epitope or an intact ligand-binding domain, nuclear localization of these mutants is being assessed in individual control transfections either by indirect immunofluorescence analysis using Mab-17 (7) or by staining with fluorescent

ER ligands (3). It is assumed that a correlation will exist between the *in vivo* potency of peptide inhibitors of ER, and their ability to accumulate in the nucleus.

Custom synthesis of estrogen receptor peptides.

We originally proposed testing the inhibitory potential of a limited number of polypeptides, focusing on the dimerization motif described by Lees *et al.* (5). Due, however, to recent advances in techniques for economical solid-phase synthesis of peptides, we have obtained a series of 40 synthetic hER peptides (Chyron Corp., La Jolla, CA), whose sequences are provided in Appendix E. Peptides # 24, 5, 42, 6, 10, and 14 overlap with the peptides proposed in the original application. Due to differences in amino acid content these peptides display a wide range of solubilities, but they all appear to be soluble in either deionized water or dimethylformamide. Dilutions (1000-fold) of each peptide stock are being tested for their ability to directly inhibit DNA-binding by hER, as described below.

In vitro gel mobility shift assay for analyzing inhibition of DNA-binding activity.

The DNA-binding activity of ER and its inhibition by synthetic peptides is being assessed using an in vitro gel mobility shift assay, essentially as described by Neff et al. (7). Briefly, ER is over expressed in Cos7 cells by transient transfection with pCMV-ER (Neff), a highly efficient expression plasmid containing a wild type hER cDNA (8). Whole cell extracts containing hER where prepared from these cells on the second day following transfection by sonicating the harvested cell pellet in a 0.4 M KCl extraction buffer containing, in addition, 20 mM HEPES (pH 7.4), 1 mM EDTA, 2 mM dithiothreitol, 10% glycerol, and 2.5 µg/ml each of antipain, aprotinin, chymostatin, and leupeptin. Following centrifugation for 10 min at 15,000 X g, aliquots of the clarified supernate (8 µl, containing 30-40 µg of protein) were preincubated for 15 min at room temperature in 20 µl of DNA-binding buffer [10 mM HEPES (pH 7.4), 1 mM MgCl<sub>2</sub>, 1 mM dithiothreitol, and 20% glycerol] also containing 1 µg tRNA, 1 µg poly(dI-dC), and 100 µg acetylated BSA. Reactions containing hER polypeptides included 2 µl of from a working dilution of the peptide stock, or approximately 20 pmol of peptide, assuming complete dissolution. Following preincubation of receptor-containing extracts in the presence or absence of the test peptide, approximately 1.2 fmol (40,000 cpm) of a <sup>32</sup>P-labeled double-stranded ERE oligonucleotide (6) were added to each sample and incubation was continued for 30 min at room temperature, followed by an additional 30 min on ice. Samples were then loaded on a preelectrophoresed non-denaturing 5% polyacrylamide gel that was run in 0.5 X TBE buffer at 280 volts for 2 hr. Gels were dried and exposed overnight for autoradiography. Inhibition of DNAbinding by ER in response to peptide addition was evidenced by a consistent decrease in the intensity of the slow-mobility hER/DNA complex. To confirm the specificity of this complex, control DNA-binding reactions were performed with equal amounts of extract from pCMV4 vectortransfected Cos7 cells and with Cos7/pCMV-ER extracts containing 1 µl of an ER-specific monoclonal antibody.

#### **RESULTS**

Specificity of DNA-binding by intact hER expressed in Cos7 cells.

A representative gel mobility shift assays is shown in Fig. 1, to demonstrate the specificity of this reaction. Intact receptor (wt hER) expressed in Cos7 cells is able to bind efficiently and with high specificity to an oligonucleotide containing a consensus estrogen response element (ERE, AGGTCACAGTGACCT) (Fig. 1, lanes 5-7), while extracts from Cos7 cells that were transfected with the pCMV4 expression vector alone (lanes 2-4) displayed only weak, nonspecific DNA-binding activity (indicated by an asterisk). The DNA-binding activity of wt hER (solid

arrow) was enhanced by pretreatment of cells expressing the receptor with 1 nM estradiol (lane 6) and was further slowed in mobility by the addition of an ER-specific monoclonal antibody, Mab-17 (lane 7), to the binding reactions (hatched arrow). It should be noted that the Mab-17 antibody failed to stabilize or alter the mobility of the non-specific DNA binding displayed by endogenous Cos7 proteins (compare lanes 3 and 4), indicating that this complex appears not to contain immunologically reactive ER. The extracts used in these binding reactions representing control and pCMV-ER transfected Cos7 cells contained comparable amounts of protein as determined using the Lowry protein assay. Since it depends on the presence of hER and is intensified by conditions that are known to stabilize DNA binding by hER, the slow mobility band (indicated by a solid arrow) can be confidently assigned to represent the hER/DNA complex.

Screening synthetic hER peptides for their ability to interfere with DNA-binding by intact receptor.

To determine whether synthetic peptides with sequences derived from the hER polypeptide chain are able to interfere with the ability of this receptor to bind efficiently to DNA, we've initiated a series of gel mobility shift assays. Each assay includes

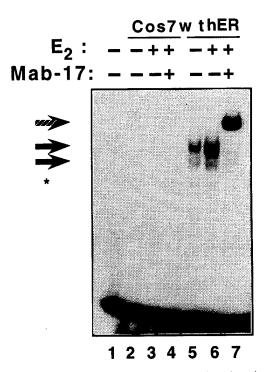


Fig. 1. Gel mobility shift assay showing the specificity of DNA complex formation by hER. Lower arrows: specific hER/DNA complex, Hatched arrow: DNA compex super-shifted by an ER-specificl antibody, asterisk: non-specific DNA complex.

control DNA-binding reactions performed with the ERE probe alone (lane 1), with protein from mock-transfected Cos7 cells (lane 2), and with protein from Cos7 cells transfected with the hER expression vector pCMV-ER (lanes 3-13). DNA-binding reactions including a single synthetic hER peptide (lanes 4-13) along with an equivalent amount of Cos7 extract were always compared with a control reaction lacking added peptide (lane 3). Representative results from two sets of experiments that have systematically screened through each of 40 synthetic peptides shown in Appendix D are given in Fig 2. As noted above, in the absence of added peptide, hER expressed in Cos7 cells is able to bind efficiently and with high specificity to an ERE-containing oligonucleotide (Fig. 2, lane 3). Our results suggest that hER suffers a dramatic reduction in DNAbinding activity in the presence of peptide P6, derived from the dimerization domain of hER (compare lane 7 with lane 3). Less dramatic inhibition of DNA-binding was caused by the addition of peptides P<sub>8</sub> (lane 9) and P<sub>10</sub> (lane 11). It is noteworthy that two of the peptides showing the greatest degree of inhibitory activity (P6 and P10) represent contiguous sequences in hER that correspond precisely to the dimerization motif identified by Parker and colleagues (5). Since each reaction included equivalent aliquots from a single hER-containing Cos7 cell extract, the intensity of the upper two hER-specific bands (indicated with solid arrows) should accurately reflect the amount of hER protein not complexed with peptide, that remains available for binding to DNA. It is also significant that a more rapidly migrating non-specific DNA complex (indicated with an asterisk) formed between an endogenous Cos7 cell protein (see lane 2) and the DNA probe shows little if any variation in intensity as a function of added peptide (lanes 2-13). Additional gel shift experiments with peptides P<sub>13</sub> through P<sub>20</sub> failed to identify other hER peptides that consistently interfered with DNA-binding, although several peptides (P<sub>16</sub>,  $P_{19}$ ,  $P_{37}$ , and  $P_{38}$ ) showed significant inhibitory activity in isolated experiments. A summary of the natural location of the peptides tested in Fig. 2 within the hER polypeptide is given in Fig. 3. Confirmation of these findings will require gel shifts such as those shown in Fig. 2 to be repeated with the entire series of synthetic hER peptides. There are also plans for the aquisition of a phosphoimage analyzer (BioRad Laboratories, Inc.) by this department that will permit the extent of DNA binding in this assay to be much more readily quantitated. Ultimately, titration experiments involving the strongest peptide inhibitors will be required to determine their relative inhibitory potency.

Dominant inhibition of the function of hER in transfected cells by expressed hER polypeptides.

A limited number of transfection experiments have been performed to test whether portions

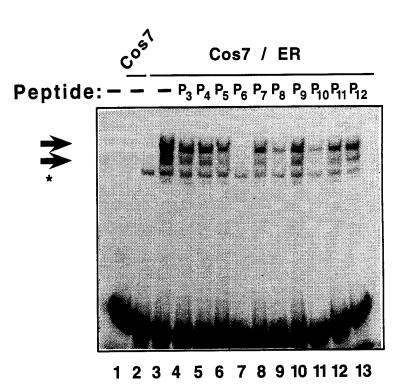


Fig. 2. Gel mobility shift assay showing inhibition of hER binding to DNA by selected synthetic hER peptides. Lane assignments are as desribed in the text. Peptide sequences are shown in Appendix E.

of the hER protein coding sequence are able to interfere with the transcriptional stimulatory activity of intact hER when they are co-expressed in cells. While it is too early to make any sweeping generalizations from these experiments, several cases have been noted where hER constructs that harbor an intact hormone binding domain possess dominant inhibitory activity, despite deletions elsewhere within the polypeptide chain. A good example of this is the repressive activity of the ERΔE3 splicing variant (6), which suffers a loss of DNA-binding ability, but retains an intact hormone binding domain. We have recently observed similar behavior for pcDNA3-ER(DEF), a construct engineered to express only the hER hormone-binding domain (residues 256-595). One possibility for explaining these results is that the presence of an exposed dimerization motif (residues 497-518) in an otherwise transcriptionally inactive receptor is sufficient to confer transdominant inhibitory activity. Transfection experiments with additional hER constructs are in progress to test this hypothesis.

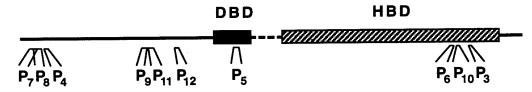


Fig. 3. Summary of the location of synthetic hER peptides used for the gel mobility shift assay shown in Fig. 2. DBD: DNA-binding domain, HBD: Hormone-binding domain.

#### **DISCUSSION**

During the first year of this award, significant progress has been made towards reaching the specific aims set out in our original application. In order to continue to pursue these research aims, a request for a change in the performance site of this project has been filed, necessitated by a change in the institutional affiliation of the PI. This is to be effected by a subcontracting arrangement under terms that have been agreed upon by the two institutions involved (The Research Foundation of SUNY and Michigan State University). A request for approval of this arrangement is pending with the US Army MRDC.

The reagents necessary for completing this project have been assembled and assays to test for the inhibitory activity of synthetic and expressed hER peptides have been developed. We have strong preliminary evidence that at least two peptides (P<sub>6</sub> and P<sub>10</sub>), and possible more, derived from the open reading frame of the hER protein appear to be capable of interfering with the DNA-binding activity of this protein. If additional gel shift experiments confirm this finding, it becomes less important to continue to employ the transient transfection assay to identify expressed hER peptides with *in vivo* dominant inhibitory activity (Task 1C). In this case, priority will be shifted away from this series of experiments. Instead, emphasis will be placed on using the *in vitro* gel mobility shift assay to define the shortest possible peptides and the optimal peptide sequence that exhibits inhibitory activity (Task 2C). Additionally, a higher priority will then be placed on testing these peptides for inhibitory activity *in vivo* using liposome-mediated peptide transfer (Task 2D).

In summary, we can report significant progress towards our goal of identifying short peptides, derived from the sequence of the hER protein, that are able to interfere with the DNA-binding activity of this receptor. The underlying justification for this work continues to be the development of a novel class of steroid receptor inhibitors that may someday prove to be useful as therapeutic agents.

## Publications During Current Award Year:

- a) Wang, Y. and Miksicek, R.J. (1994) Characterization of estrogen receptor cDNAs from human uterus: identification of a novel *Pvu II* polymorphism. *Mol. Cell Endocrinol.* 101:101-110.
- b) Miksicek, R.J. (1994) Steroid receptor variants and their potential role in cancer. Seminars in Cancer Biology 5:369-379.
- c) Elias, J., Hyder, D.M., Miksicek, R.J., Heimann, A., and Marigiota, M. (1995) Interpretation of steroid receptors in breast cancer: Report of a case with discordant estrogen receptor results using ER1D5 and H222 antibodies. *J Histotechnology* 18(4), in press
- d) Miksicek, R.J., Carlson, K.E., Hwang, K.-J., and Katzenellenbogen, J.A. (1995) Studies using fluorescent tetrahydrochrysene estrogens for *in situ* visualization of the estrogen receptor in living cells. *Mol. Endocrinol.* 9:592-604.

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Miksicek, R.J., Carlson, K.E., Hwang, K.-J., and Katzenellenbogen, J.A. (1995) Studies using fluorescent tetrahydrochrysene estrogens for *in situ* visualization of the estrogen receptor in living cells. *Mol. Endocrinol.* 9:592-604.

Tora, L., White, J., Brou, C., Tasset, D., Webster, N., Scheer, E., Chambon, P. (1989) The human estrogen receptor has two independent nonacidic transcriptional activation functions *Cell* 

**59**:477-487.

- Lees, J.A., Fawell, S.E., White, R., Parker, M.G. (1990) A 22-amino acid peptide restores DNA-binding activity to dimerization defective mutants of the estrogen receptor. *Mol. Cell. Biol.* 10:5529-663.
- Wang, Y. and Miksicek, R.J. (1991) Identification of a dominant negative form of the human estrogen receptor. *Mol. Endocrinol.* 5:1707-1715.
- 7) Neff, S., Sadowski, C., and Miksicek, R.J. (1993) Site-directed mutagenesis of the hormone-binding domain of the human estrogen receptor reveals that two cysteine residues play a role in DNA-binding and subcellular localization. *Mol. Endocrinol.* 8:1215-1223.
- 8) Wang, Y. and Miksicek, R.J. (1994) Characterization of estrogen receptor cDNAs from human uterus: identification of a novel *Pvu II* polymorphism. *Mol. Cell. Endocrinol.* 112:1-13.

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# Appendix A: Letters of Appointment for PI

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Office of the Vice President for Health Sciences

June 15, 1995

Richard Miksicek, Ph.D. 14 Paul Street Port Jefferson Station, NY 11776

Dear Dr. Miksicek:

On behalf of President Kenny, I am writing to confirm the end of your temporary salaried appointment as Research Assistant Professor in the Department of Pharmacological Sciences in the School of Medicine effective June 30, 1995.

In view of your desire to continue your affiliation with the State University of New York at Stony Brook, effective July 1, 1995, I am pleased to grant you a non-salaried appointment as Research Assistant Professor as a member of the voluntary faculty in the in the School of Medicine. This appointment is at the pleasure of the President and subject to the <u>Policies</u> of the Board of Trustees governing such appointments.

If you wish to accept this appointment, please sign and date the endorsement on the enclosed copy of this letter and return that copy to me by JUly 15, 1995.

Sincerely

H. Oaks

Vice President for Health Sciences

JHO:ha

cc: Dr. B. Hool

Dr. N. N. Abumrad

Dr. A. P. Grollman

State University of New York at Stony Brook Stony Brook, New York 11794-8400

516-444-2101 Fax: 516-444-2509

DAMD 1794J4411 MIKSICEK, Richard J.

## Appendix B: Letter Requesting Approval for Subcontract Agreement

M.S.U. C.G.A.

ID:5173539812

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11:58 No.003 P.02

# MICHIGAN STATE

June 13, 1995

Mr. Mike Younkins Grants and Contracts Officer US Army Medical Research and Development Command Fort Detrick, MD 21702-5012

re:

Breast Cancer Research Award

PI: Richard J. Miksicek

Title: Development of Novel Peptide Inhibitors of the Estrogen

Receptor

Award Period:

10/94-09/96

Code: A125

ID: DAMD-17-12J-4411

#### Dear Mr. Younkins:

In relation to the prior request of Dr. Richard Miksicek (in a letter to you dated May 22, 1995), arrangements are being made as of July 1, 1995, to move the above referenced US Army MRDC Breast Cancer Research Project to the Department of Physiology, Michigan State University. In consultation with Dr. Miksicek and the grants and contracts officers of our respective institutions, I will be assuming the role of Principal Investigator on the balance of the research award that is to be subcontracted to Michigan State University. I currently hold an appointment as Professor of Physiology as well as Director of the Breast Cancer Program of the Michigan State University Cancer Center. My qualifications to assume the administrative and supervisory role on this project include the prior and current direction of a number of federally sponsored research projects in the area of hormonal control of the development and physiology of the breast. In addition, Dr. Miksicek will maintain a close supervisory role during the duration of the project period.

With this letter I am requesting that the Office of Contracts and Grants Administration of Michigan State University be permitted to initiate negotiations with the Research Foundation of SUNY to subcontract the unobligated balance of this research contract to Michigan State University for the duration of the award period (07/01/95 through 09/30/98). All terms of the parent contract, if acceptable to Michigan State University, including the specific aims, the award period, and the total award amount will remain unchanged, although some rebudgeting of funds between categories will occur as part of the subcontract. A revised budget is being prepared in consultation with the grants and contracts administrators of the respective institutions. This budget will be submitted to you for approval as soon as it is available.



DEPARTMENT OF PHYSIOLOGY

Michigan State University 108 Giltner Hall East Lansing, Michigan 48824-1101 517/355-6475 FAX: 517/355-5125 M.S.U. C.G.A.

ID:5173539812

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Mr. Younkins June 13, 1995 Page 2

The names and phone numbers of the administrators at Stony Brook and MSU are as follows:

Michigan State University:

Suzanne Morgan
Contract and Grant Administration
Michigan State University
East Lansing, MI 48824
Phone: (517)-353-7297
FAX: (517)-353-9812

email: CGAO6@msu.edu

SUNY at Stony Brook:

Barbara Harris, Associate for Sponsored Research The Research Foundation of SUNY Office of Research Services State University of New York at Stony Brook Stony Brook, NY 11794-3366 Phone: (516) 632 9038

Phone: (516)-632-9028 FAX: (516)-632-6963

email: bharris@ors.ors.sunysb.edu

I would greatly appreciate a letter from you providing authorization to proceed with this arrangement. Also please appraise me of any certifications or approvals that will be required by the US Army MRDC from myself or the Grants and Contracts Office of Michigan State University in conjunction with the subcontract. Thank you for your prompt attention to this matter.

Sincerely yours,

Sandra Haslam, Ph.D. Professor of Physiology

Approved:

William S. Spielman, Ph.D. Professor and Chairperson

Daniel Evon Assistant Director

Contract and Grant Administration

SHjr

## Appendix C: Revised Description of Facilities and Equipment

Permission has been requested to move the performance site of this project as of 07/01/95 from SUNY at Stony Brook to the Department of Physiology at Michigan State University (MSU). To effect this move, we are proposing that the parent award will remain with the Research Foundation of the State University of New York and, as agreed upon by both institutions, unobligated costs will be subcontracted to MSU.

The laboratory assigned to Dr. Miksicek at MSU occupies 850 square feet of space on the 2nd floor of Giltner Hall on the MSU campus in East Lansing, MI. It is fully equipped for the proposed biochemical and molecular biological research. The laboratory contains a separate area for tissue culture work. Cold room and darkroom facilities are available on site.

Items of equipment available for exclusive use of Dr. Miksicek and his associates include a Pharmacia Ultrospec II spectrophotometer, electrophoresis chambers and power supplies, standard and low temperature freezers, refrigerated centrifuges, microcentrifuges, a Savant Speed Vac, Savant thermal cycler, balances, a pH meter, and a Millipore water purification system. A designated cell culture room adjacent to the research lab houses a laminar flow hood, a humidified CO<sub>2</sub> incubator, a Nikon inverted microscope, and a table top centrifuge. The following equipment is shared with other departmental faculty located in Giltner Hall: ultracentrifuges, liquid scintillation counters, a spectrofluorimeter, a Perkin-Elmer thermal cycler, low temperature freezers, a Zeiss photomicroscope equipped for epifluorescence illumination, dark room and X-ray film developing equipment.

Personnel have access to personal computers which include an ethernet connection to internet. Programs for DNA sequence analysis are available as a GCG software package. All investigators involved with this research have access to relevant databases, literature retrieval (Medline) services, and computer networks linking them to other universities and research institutions. Office space (100 sq ft) is assigned to the Co-Principal Investigator immediately adjacent to the laboratory space. Individual desks have been made available to all research personal adjacent to their work areas. Two personal computers for word processing and data management are present for the sole use of this research staff.

Access to custom oligonucleotide synthesis is available through the Depts. of Microbiology and Biochemistry on a cost-for-service basis. The university operates machine and electronic shops that are available to faculty.

Arrangements for disposal and storage of radioactive wastes are provided as an institutional service by the university. Dr. Miksicek is fully licensed and registered to use radioisotopes on the MSU campus.

## Appendix D: Sequences of Synthetic Peptides Obtained for this Project

\* note, these peptides are numbered 3-42 based on increasing size (peptides 1 and 2 were synthesis controls), but they have been ordered by their position within the protein sequence or ER, beginning at the N-terminus.

Number		<u>Peptide Sequence</u>	i	Size (aa residues)
37)	free NH <sub>2</sub> -	MTMTLHTKASGMALLHQIQC	-amide	20
15)	acetyl-	GNELEPLNRPQLKI	-amide	14
7)	acetyl-	PLERPLGEVYL	-amide	11
8)	acetyl-	PLGEVYLDSSK	-amide	11
4)	acetyl-	PAVYNYPEG	-amide	9
38)	acetyl-	AAYEFNAAAAANAQVYGQTG	-amide	20
32)	acetyl-	LPYGPGSEAAAFGSNGLG	-amide	18
33)	acetyl-	GFPPLNSVSPSPLMLLHP	-amide	18
9)	acetyl-	PPQLSPFLQPHG	-amide	12
11)	acetyl-	QQVPYYLENEPSG	-amide	13
12)	acetyl-	YTVREAGPPAFYR	-amide	13
29)	acetyl-	PNSDNRRQGGRERLASC	-amide	17
23)	acetyl-	NDKGSMAMESAKETRY	-amide	16
16)	acetyl-	CNDYASGYHYGVWS	-amide	14
24)	acetyl-	CKAFFKRSIQGHNDYM	-amide	16
5)	acetyl-	CTIDKNRRKS	-amide	10
39)	acetyl-	CYEVGMMKGGIRKDRRGGRM	-amide	20
40)	acetyl-	LKHKRQRDDGEGRGEVGSAG	-amide	20
41)	acetyl-	DMRAANLWPSPLMIKRSKKN	-amide	20
36)	acetyl-	SLALSLTADQMVSALLDAE	-amide	19
13)	acetyl-	PPILYSEYDPTRP	-amide	13
34)	acetyl-	FSEASMMGLLTNLADREL	-amide	18
25)	acetyl-	VHMINWAKRVPGFVDL	-amide	16
17)	acetyl-	TLHDQVHLLECAWL	-amide	14
18)	acetyl-	EILMIGLVWRSMEH	-amide	14
26)	acetyl-	PGKLLFAPNLLLDRNQ	-amide	16
30)	acetyl-	GKCVEGMVEIFDMLLAT	-amide	17
27)	acetyl-	SSRFRMMNLQGEEFVC	-amide	16
28)	acetyl-	LKSIILLNSGVYTFLS	-amide	16
35)	acetyl-	STLKSLEEKDHIHRVLDK	-amide	18
22)	acetyl-	ITDTLIHLMAKAGLT	-amide	15
42)	acetyl-	LQQQHQRLAQLLLILSHIRH	-amide	20
6)	acetyl-	LQQQHQRLAQ	-amide	10
10)	acetyl-	LLLILSHIRHMS	-amide	12
14)	acetyl-	MSNKGMEHLYSMK	-amide	13
3)	acetyl-	CKNVVPLY	-amide	8
19)	acetyl-	DLLLEMLDAHRLHA	-amide	14
20)	acetyl-	PTSRGGASVEETDQ	-amide	20
21)	acetyl-	SHLATAGSTSSHSL	-amide	20
31)	acetyl-	CQKYYITGEAEGFPATV	-free a	cid 17